



XML: Bioinformatics

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XML for Bioinformatics Ethan Cerami, 2006-06-02 Introduction The goal of this book is to introduce XML to a bioinformatics audience It does so by introducing the fundamentals of XML Document Type Definitions DTDs XML Namespaces XML Schema and XML parsing and illustrating these concepts with specific bioinformatics case studies The book does not assume any previous knowledge of XML and is geared toward those who want a solid introduction to fundamental XML concepts The book is divided into nine chapters Chapter 1 Introduction to XML for Bioinformatics This chapter provides an introduction to XML and describes the use of XML in biological data exchange A bird's eye view of our first case study the Distributed Annotation System DAS is provided and we examine a sample DAS XML document The chapter concludes with a discussion of the pros and cons of using XML in bioinformatic applications Chapter 2 Fundamentals of XML and BSML This chapter introduces the fundamental concepts of XML and the Bioinformatic Sequence Markup Language BSML We explore the origins of XML define basic rules for XML document structure and introduce XML Namespaces We also explore several sample BSML documents and visualize these documents in the TM Rescentris Genomic Workspace Viewer **Xml For**

Bioinformatics Cerami, 2005-01-01 *Biological and Medical Data Analysis* José Luis Oliveira, 2005-11-04 This book constitutes the refereed proceedings of the 6th International Symposium on Biological and Medical Data Analysis ISBMDA 2005 held in Aveiro Portugal in November 2005 The 39 revised full papers presented were carefully reviewed and selected for inclusion in the book The papers are organized in topical sections on medical databases and information systems data analysis and image processing knowledge discovery and data mining statistical methods and tools for biomedical data analysis decision support systems collaborative systems in biomedical informatics as well as computational models structural analysis and microarray data analysis in the scope of bioinformatics Current Trends in Database Technology - EDBT 2006

Torsten Grust, Hagen Höpfner, Arantza Illarramendi, Stefan Jablonski, Marco Mesiti, Sascha Müller, Paula-Lavinia Patranjan, Kai-Uwe Sattler, Myra Spiliopoulou, Jef Wijsen, 2006-10-17 This book constitutes the thoroughly refereed joint post proceedings of nine workshops held as part of the 10th International Conference on Extending Database Technology EDBT 2006 held in Munich Germany in March 2006 The 70 revised full papers presented were selected from numerous submissions during two rounds of reviewing and revision *Bioinformatics in the Post-genomic Era* Jeffrey Augen, 2005 A comprehensive treatment of the role of bioinformatics in the emerging world of molecular medicine for anyone involved in this new field

The Grid Maozhen Li, Mark Baker, 2005-11-01 Find out which technologies enable the Grid and how to employ them successfully This invaluable text provides a complete clear systematic and practical understanding of the technologies that enable the Grid The authors outline all the components necessary to create a Grid infrastructure that enables support for a range of wide area distributed applications The Grid Core Technologies takes a pragmatic approach with numerous practical examples of software in context It describes the middleware components of the Grid step by step and gives hands on advice

on designing and building a Grid environment with the Globus Toolkit as well as writing applications The Grid Core Technologies Provides a solid and up to date introduction to the technologies that underpin the Grid Contains a systematic explanation of the Grid including its infrastructure basic services job management user interaction and applications Explains in detail OGSA Open Grid Services Architecture Web Services technologies SOAP WSDL UDDI and Grid Monitoring Covers Web portal based tools such as the Java CoG GridPort GridSphere and JSR 168 Portlets Tackles hot topics such as WSRF Web Services Resource Framework the Semantic Grid the Grid Security Infrastructure and Workflow systems Offers practical examples to enhance the understanding and use of Grid components and the associated tools This rich resource will be essential reading for researchers and postgraduate students in computing and engineering departments IT professionals in distributed computing as well as Grid end users such as physicists statisticians biologists and chemists

Ontologies for Bioinformatics Kenneth Baclawski, Tianhua Niu, 2006 Ontologies as a critical framework for the vast amounts of data in the postgenomic era an introduction to the basic concepts and applications of ontologies and ontology languages for the life sciences Recent advances in biotechnology spurred by the Human Genome Project have resulted in the accumulation of vast amounts of new data Ontologies computer readable precise formulations of concepts and the relationship among them in a given field are a critical framework for coping with the exponential growth of valuable biological data generated by high output technologies This book introduces the key concepts and applications of ontologies and ontology languages in bioinformatics and will be an essential guide for bioinformaticists computer scientists and life science researchers The three parts of Ontologies for Bioinformatics ask and answer three pivotal questions what ontologies are how ontologies are used and what ontologies could be which focuses on how ontologies could be used for reasoning with uncertainty The authors first introduce the notion of an ontology from hierarchically organized ontologies to more general network organizations and survey the best known ontologies in biology and medicine They show how to construct and use ontologies classifying uses into three categories querying viewing and transforming data to serve diverse purposes Contrasting deductive or Boolean logic with inductive reasoning they describe the goal of a synthesis that supports both styles of reasoning They discuss Bayesian networks as a way of expressing uncertainty describe data fusion and propose that the World Wide Web can be extended to support reasoning with uncertainty They call this inductive reasoning web the Bayesian web Medinfo 2007

Klaus A. Kuhn, James R. Warren, Tze-Yun Leong, 2007 The papers presented are refereed and from all over the world They reflect the breadth and depth of the field of biomedical and health informatics covering topics such as health information systems knowledge and data management education standards consumer health and human factors emerging technologies sustainability organizational and economic issues genomics and image and signal processing As this volume carries such a wide collection it will be of great interest to anyone engaged in biomedical and health informatics research and application

XML and Web Services Unleashed Ron Schmelzer, 2002 The complete and up to date XML reference book on topics that

matter The XMLiterate developer is certain to find plenty of new topics and emerging specs that have not been explored previously

Database and Expert Systems Applications Hendrik Decker, Lenka Lhotská, Sebastian Link, Marcus Spies, Roland R. Wagner, 2014-08-20 This two volume set LNCS 8644 and LNCS 8645 constitutes the refereed proceedings of the 25th International Conference on Database and Expert Systems Applications DEXA 2014 held in Munich Germany September 1-4 2014 The 37 revised full papers presented together with 46 short papers and 2 keynote talks were carefully reviewed and selected from 159 submissions The papers discuss a range of topics including data quality social web XML keyword search skyline queries graph algorithms information retrieval XML security semantic web classification and clustering queries social computing similarity search ranking data mining big data approximations privacy data exchange data integration web semantics repositories partitioning and business applications *XML Data Representation and Transformations for Bioinformatics* Kevin C. Woods, California State University, Sacramento, 2007

Biological and Medical Data Analysis, 2005 **IBM Systems Journal** International Business Machines Corporation, 2007 **Genome Research**, 2003 **Proceedings of the National Conference on Computing for Nation Development**, 2007 *IBM DB2 9 New Features* Paul Zikopoulos, George Baklarz, Leon Katsnelson, Chris Eaton, 2007-03-05 Design and deploy integrated enterprise class business solutions Enhance productivity profitability and efficiency using IBM DB2 9 and the detailed information in this comprehensive resource Written by experienced IBM data server professionals IBM DB2 9 New Features covers the technical details of the major new aspects of DB2 9 alongside real world business uses syntax examples and insider tips Learn how to manipulate and manage XML data using pureXML automate maintenance tasks and seamlessly integrate Java PHP and Perl scripts Plus you'll get vital coverage of the brand new DB2 Developer Workbench enhanced NET developer support Ruby on Rails and database security Learn about all the auto configuration features that make getting DB2 up and running faster than ever before Define share and validate data using XML and the unique pureXML DB2 Hybrid Storage Engine Improve runtime performance and optimize storage with the new DB2 Compression Technology Automatically handle DB2 memory allocation using the Self Tuning Memory Manager Use Automatic Storage Autonomic Computing and a host of other autonomic features that make managing DB2 a snap Create dynamic data server blueprints with Rational Data Architect version 6.1 Incorporate Java applications using the IBM WebSphere Application Server and DB2 data server Develop open source scripts using PHP Perl Toad and Ruby on Rails Utilize the DB2 Developer Workbench or exploit the enhanced NET developer support Deploy the latest DB2 security and high availability features *Very Large Data Bases*, 2005 **VLDB 2005**, 2005 **Dr. Dobb's Journal**, 2006 *Data Integration in the Life Sciences*, 2005

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